NUCLEOTIDE AND AA SEQUENCE

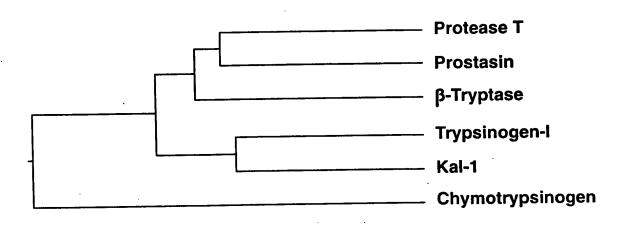
Nucleic acid sequence of Protease T (SEQ.ID.NO.:1)

GACCACGCCCTGCGCCCCAGCCAGGCCTGAGGACATGAGGCGGCGGCGGCGGCGGTGCCGC TCCTGCTGCTGTGTTTTGGGTCTCAGAGGGCCAAGGCAGCAACAGCCTGTGGTCGCC CCAGGATGCTGAACCGAATGGTGGGCGGGCAGGACACGCAGGAGGGCGAGTGGCCCTGGC AAGTCAGCATCCAGCGCAACGGAAGCCACTTCTGCGGGGGCAGCCTCATCGCGGAGCAGT GGGTCCTGACGGCTGCGCACTGCTTCCGCAACACCTCTGAGACGTCCCTGTACCAGGTCC AGGTGGAGAGCAACCCCCTGTACCAGGGCACGGCCTCCAGCGCTGACGTGGCCCTGGTGG AAGACCTCCTGCCCGAACCGCGGATCCTGCAGAAACTCGCTGTGCCCCATCATCGACACAC CCAAGIGCAACCTGCTCTACAGCAAAGACACCGAGTTTGGCTACCAACCCAAAACCATCA AGAATGACATGCTGTGCGCCGGCTTCGAGGGGGCAAGAAGGATGCCTGCAAGGGCGACT CGGGGGGCCCCTGGTGTGCCTCGTGGGTCAGTCGTGGCTGCAGGCGGGGGTGATCAGCT GGGGTGAGGGCTGTGCCCGCCAGAACCGCCCAGGTGTCTACATCCGTGTCACCGCCCACC AGAAGTGAGACCCCCGGGGCCAGGAGCCCCTTGAGCAGAGCTCTGCACCCAGCCTGCCCG CCCACACCATCCTGCTGCTCCCCAGCGCTGCTGTTGCACCTGTGAGCCCCACCAGACT CATTTGTAAATAGCGCTCCTTCCTCCCCCTCTCAAATACCCTTATTTTTATTTTATGTTTCTC CCAATAAAAACCCAGCCTGTGTGCCAGCTG

>Amino Acid Sequence of Protease T (SEQ.ID.NO.:7)

MRRPAAVPILILICFGSQRAKAATACGRPRMLNRMVGGQDTQEGEWPWQVSIQRNGSHFC GGSLIAEQWVLTAAHCFRNTSETSLYQVLLGARQLVQPGPHAMYARVRQVESNPLYQGTA SSADVALVELEAPVPFTNYILPVCLPDPSVIFETGMVCWVTGWGSPSEEDLLPEPRILQK LAVPIIDTPKCNLLYSKDTEFGYQPKTIKNDMLCAGFEEGKKDACKGDSGGPLVCLVGQS WLQAGVISWGEGCARQNRPGVYIRVTAHHNWIHRIIPKLQFQPARLGGQK

FIG. 2



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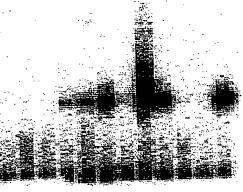
Prot.

<u>م:</u>

LIVER LUNG LYMPH NODE MONOCYTE (PBM) OSTEOSARCOMA OVARY PANCREAS PITUITARY PLACENTA PROSTATE RETINA SKELETAL MUSCLE

BCELL LYMPHOMA
BONE MARROW
AMYGDALA
CAUDATE NUCLEUS
CEREBELLUM
CEREBELLUM
CEREBRAL CORTEX
HIPPOCAMPUS
HYPOTHALAMUS
SUBSTANTIA NIGRA
SUBTHALMIC NUCLEUS
THALAMUS
BREAST CARCINOMA





COLON ENDOTHELIAL EPIDERMIS FAT CELL FETAL BRAIN FIBROBLAST HEART JURKAT JURKAT+PMA+PHA-M KERATINOCYTE KIDNEY

3/6

FUSION PROTEIN

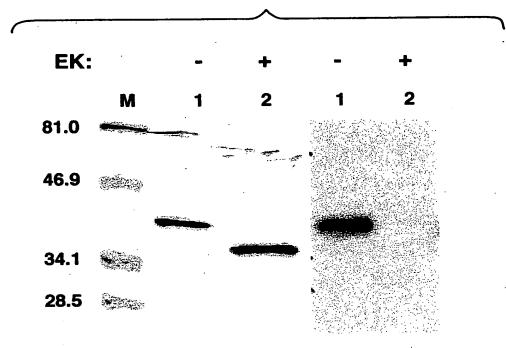
Nucleic Acid sequence of PFEK-PROTT-HIS fusion protein (SEQ.ID.NO.:8).

GAATTCACCACCATGGACAGCAAAGGTTCGTCGCAGAAATCCCGCCTGCTCCTGCTGCTG GTGGTGTCAAATCTACTCTTGTGCCAGGGTGTGGTCTCCGACTACAAGGACGACGACGAC GTGGACGCGCCCCTTTGCTGCCCCCTTTGATGATGATGACAAGATCGTTGGGGGCTAT GCTCTAGAGGAGGGCGAGTGGCCCTGGCAAGTCAGCATCCAGCGCAACGGAAGCCACTTC TGCGGGGCAGCTCATCGCGGAGCAGTGGGTCCTGACGGCTGCGCACTGCTTCCGCAAC ACCTCTGAGACGTCCCTGTACCAGGTCCTGCTGGGGGGCAAGGCAGCTAGTGCAGCCGGGA CCACACGCTATGTATGCCCGGGTGAGGCAGGTGGAGAGCAACCCCCTGTACCAGGGCACG GCCTCCAGCGCTGACGTGGCCCTGGTGGAGCTGGAGCACCAGTGCCCTTCACCAATTAC ATCCTCCCGTGTGCCTGACCCCTCGGTGATCTTTGAGACGGCATGAACTGCTGG GTCACTGGCTGGGGCAGCCCCAGTGAGGAAGACCTCCTGCCCGAACCGCGGATCCTGCAG AAACTCGCTGTGCCCATCATCGACACACCCAAGTGCAACCTGCTCTACAGCAAAGACACC GAGTTTGGCTACCAACCCAAAACCATCAAGAATGACATGCTGTGCGCCGGCTTCGAGGAG GGCAAGAAGGATGCCTGCAAGGGCGACTCGGGCGCCCCCTGGTGTGCCTCGTGGGTCAG TCGTGGCTGCAGGCGGGGTGATCAGCTGGGGTGAGGGCTGTGCCCGCCAGAACCGCCCA GGTGTCTACATCCGTGTCACCGCCCACCACAACTGGATCCATCGGATCATCCCCAAACTG CAGTTCCAGCCAGCGAGGTTGGGCGGCCAGAAGTCTAGACATCACCATCACCATCACTAG CGGCCGCTTCCCTTTAGTGAGGGTTAATGCTTCGAGCAGACATGATAAGATACATTGATG AGTTTGGACAAACCACAACTAGAATGCAGTGAAAAAAATGCTTTATTTGTGAAATTTGTG ATGCTATTGCTTTATTTGTAACCATTATAAGCTGCAATAAACAAGTTGAC

Amino Acid sequence of PFEK-PROTT-HIS fusion protein (SEQ.ID.NO.:9).

MDSKGSSQKSRLLLLLVVSNLLLCQGVVSDYKDDDDVDAAALAAPFDDDDKIVGGYALEE GEWPWQVSIQRNGSHFCGGSLIAEQWVLTAAHCFRNTSETSLYQVLLGARQLVQPGPHAM YARVRQVESNPLYQGTASSADVALVELEAPVPFTNYILPVCLPDPSVIFETGMNCWVTGW GSPSEEDLLPEPRILQKLAVPIIDTPKCNLLYSKDTEFGYQPKTIKNDMLCAGFEEGKKD ACKGDSGGPLVCLVGQSWLQAGVISWGEGCARQNRPGVYIRVTAHHNWIHRIIPKLQFQP ARLGGQKSRHHHHHH





Protease: PFEK-Protease T-HIS

Protease Activity of Protease T

H-D-Pro-HHT-Arg-pNA
H-D-Lys(CBO)-Pro-Arg-pNA
H-DL-Val-Leu-Arg-pNA

